

# Package ‘modelfree’

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**Title** Model-free estimation of a psychometric function

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**Depends** R (>= 2.8.1), PolynomF (>= 0.93), SparseM (>= 0.79), stats,utils, base

**LazyLoad** yes

**LazyData** yes

**Description** Local linear estimation of psychometric functions. Provides functions for nonparametric estimation of a psychometric function and for estimation of a derived threshold and slope, and their standard deviations and confidence intervals

**URL** <http://www.modelfree.manchester.ac.uk>, <http://www.modelfree.liv.ac.uk>,  
<http://modelfree.r-forge.r-project.org/>

**License** GPL (>= 2)

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01_Miranda	<i>Example 1. Frequency of seeing</i>
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### **Description**

A flash of light of variable intensity was presented repeatedly at a fixed location in the visual field of a subject who reported whether the flash was visible.

### **Usage**

```
data( "01_Miranda" )
```

### **Format**

Data frame with three columns for 10 stimulus levels: x: Stimulus level. r: Number of successes. m: Number of trials.

### **References**

Miranda, M. A. & Henson, D. B. "Perimetric sensitivity and response variability in glaucoma with single-stimulus automated perimetry and multiple-stimulus perimetry with verbal feedback", *Acta Ophthalmologica*, 86, 202-206, 2008.

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02\_Levi*Example 2. Visual detection of path deviation*

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**Description**

The subject was presented with the image of a dot moving rightwards on a linear path until it reached the midline of the display, when it changed direction either upwards or downwards. The subject had to indicate the direction.

**Usage**

```
data( "02_Levi" )
```

**Format**

Data frame with three columns for 7 stimulus levels: x: Stimulus level. r: Number of successes. m: Number of trials.

**References**

Levi, D. M. & Tripathy, S. P. "Is the ability to identify deviations in multiple trajectories compromised by amblyopia?", *Journal of Vision*, 6(12), 1367-1379, 2006.

---

03\_Carcagno*Example 3. Discrimination of pitch*

---

**Description**

The subject had to identify the interval containing a tone whose fundamental frequency was different from that in the other two intervals.

**Usage**

```
data( "03_Carcagno" )
```

**Format**

Data frame with three columns for 8 stimulus levels: x: Stimulus level. r: Number of successes. m: Number of trials.

**References**

Unpublished data from S. Carcagno, Lancaster University, July 2008

---

04\_Xie*Example 4. Discrimination of 'porthole' views of natural scenes*

---

**Description**

The subject was presented with a display split into two parts, one containing a pair of patches from the same image, the other a pair from different images, and the subject had to judge which pair came from the same image.

**Usage**

```
data( "04_Xie" )
```

**Format**

Data frame with three columns for 10 stimulus levels: x: Stimulus level. r: Number of successes. m: Number of trials.

**References**

Xie, Y. \& Griffin, L. D. "A 'portholes' experiment for probing perception of small patches of natural images", *Perception*, 36, 315, 2007.

---

05\_Schofield*Example 5. Induction of a visual motion aftereffect*

---

**Description**

The subject was presented with a moving adaptation stimulus, followed by a test stimulus.

**Usage**

```
data( "05_Schofield" )
```

**Format**

Data frame with three columns for 7 stimulus levels: x: Stimulus level. r: Number of successes. m: Number of trials.

**References**

Schofield, A. J., Ledgeway, T., \& Hutchinson, C. V. "Asymmetric transfer of the dynamic motion aftereffect between first- and second-order cues and among different second-order cues", *Journal of Vision*, 7(8), 1-12, 2007.

---

06\_Nascimento*Example 6. Discrimination of image approximations*

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**Description**

The subject was shown an image of a natural scene and an approximation of this image based on principal component analysis. The task was to distinguish between the images.

**Usage**

```
data( "06_Nascimento" )
```

**Format**

Data frame with three columns for 8 stimulus levels: x: Stimulus level. r: Number of successes. m: Number of trials.

**References**

Nascimento, S.M.C., Foster, D.H., \& Amano, K. "Psychophysical estimates of the number of spectral-reflectance basis functions needed to reproduce natural scenes", Journal of the Optical Society of America A-Optics Image Science and Vision, 22 (6), 1017-1022, 2005.

---

07\_Baker*Example 7. Auditory detection of a gap in noise*

---

**Description**

A 300-ms noise burst containing a gap of 2–8 ms duration or no gap was presented to one ear of a subject.

**Usage**

```
data( "07_Baker" )
```

**Format**

Data frame with three columns for 8 stimulus levels: x: Stimulus level. r: Number of successes. m: Number of trials.

**References**

Baker, R. J., Jayewardene, D., Sayle, C., \& Saeed, S. "Failure to find asymmetry in auditory gap detection", Laterality: Asymmetries of Body, Brain and Cognition, 13, 1-21, 2008.

---

bandwidth_bootstrap	<i>Bootstrap estimate of bandwidth</i>
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### Description

Finds bootstrap estimate of the optimal bandwidth  $h$  for binomial data in local polynomial estimation of psychometric function (PF) with guessing and lapsing rates specified in `lims`.

### Usage

```
bandwidth_bootstrap( r, m, x, H, N, h0 = NULL, link = c( "logit" ), guessing = 0, lapsing = 0, K
```

### Arguments

<code>r</code>	number of successes in points <code>x</code>
<code>m</code>	number of trials in points <code>x</code>
<code>x</code>	stimulus levels
<code>H</code>	minimum and maximum values of bandwidth to be considered
<code>N</code>	number of bootstrap replications
<code>h0</code>	pilot bandwidth; if not specified, then the scaled plug-in bandwidth is used
<code>link</code>	name of the link function to be used; default is "logit"
<code>guessing</code>	guessing rate; default is 0
<code>lapsing</code>	lapsing rate; default is 0
<code>K</code>	power parameter for Weibull and reverse Weibull link; default is 2
<code>p</code>	order of the polynomial; default is 1
<code>ker</code>	kernel function for weights; default "dnorm"
<code>maxiter</code>	maximum number of iterations in Fisher scoring; default is 50
<code>tol</code>	tolerance level at which to stop Fisher scoring; default is 1e-6
<code>method</code>	loss function to be used in bandwidth: choose from: 'ISEeta', 'ISE', 'deviance'; default is "all"

### Value

<code>h</code>	bootstrap bandwidth for the chosen method; if no method was specified, then it a list of three elements with entries corresponding to the estimated bandwidths on p-scale ( <code>h\$pscale</code> ), on eta-scale ( <code>h\$etascale</code> ) and for mean likelihood ( <code>h\$deviance</code> )
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### Examples

```
data("01_Miranda")
h<- bandwidth_bootstrap( example01$r, example01$m, example01$x, c( 0.1, 10 ), 10 )
```

---

bandwidth\_cross\_validation

*Cross-validation estimate of bandwidth*


---

## Description

Finds the cross-validation bandwidth for the local polynomial estimator of the psychometric function (PF) with guessing and lapsing rates specified in `lims`.

## Usage

```
bandwidth_cross_validation( r, m, x, H, link = c( "logit" ), guessing = 0, lapsing = 0, K = 2, p
```

## Arguments

<code>r</code>	number of successes in points <code>x</code>
<code>m</code>	number of trials in points <code>x</code>
<code>x</code>	stimulus levels
<code>H</code>	minimum and maximum values of bandwidth to be considered
<code>link</code>	name of the link function to be used; default is "logit"
<code>guessing</code>	guessing rate; default is 0
<code>lapsing</code>	lapsing rate; default is 0
<code>K</code>	power parameter for Weibull and reverse Weibull link; default is 2
<code>p</code>	order of the polynomial; default is 1
<code>ker</code>	kernel function for weights; default "dnorm"
<code>maxiter</code>	maximum number of iterations in Fisher scoring; default is 50
<code>tol</code>	tolerance level at which to stop Fisher scoring; default is 1e-6
<code>method</code>	loss function to be used in cross-validation: choose from: 'ISEeta', 'ISE', 'deviance'; default is "all"

## Value

<code>h</code>	cross-validation bandwidth for the chosen method; if no method was specified, then it a list of three elements with entries corresponding to the estimated bandwidths on p-scale ( <code>h\$pscale</code> ), on eta-scale ( <code>h\$etascale</code> ) and for mean likelihood ( <code>h\$deviance</code> )
----------------	---

## Examples

```
data("01_Miranda")
h<- bandwidth_cross_validation( example01$r, example01$m, example01$x, c( 0.1, 10 ) )
```

---

bandwidth_optimal	<i>Cross-validation estimate of bandwidth for known distributions</i>
-------------------	---

---

### Description

Finds the cross-validation bandwidth for the local polynomial estimator of the psychometric function (PF) with guessing and lapsing rates specified in `lims`. The difference between this function and `bandwidth_cross_validation` is that here the true psychometric function is known.

### Usage

```
bandwidth_optimal( ptrue, r, m, x, H, link = c( "logit" ), guessing = 0, lapsing = 0, K = 2, p =
```

### Arguments

<code>ptrue</code>	the true function. Vector with the value of PF at each design point
<code>r</code>	number of successes in points <code>x</code>
<code>m</code>	number of trials in points <code>x</code>
<code>x</code>	design points
<code>H</code>	minimum and maximum values of bandwidth to be considered
<code>link</code>	name of the link function to be used; default is "logit"
<code>guessing</code>	guessing rate; default is 0
<code>lapsing</code>	lapsing rate; default is 0
<code>K</code>	power parameter for Weibull and reverse Weibull link; default is 2
<code>p</code>	order of the polynomial; default is 1
<code>ker</code>	kernel function for weights; default "dnorm"
<code>maxiter</code>	maximum number of iterations in Fisher scoring; default is 50
<code>tol</code>	tolerance level at which to stop Fisher scoring; default is 1e-6
<code>method</code>	loss function to be used in cross-validation: choose from: 'ISEeta', 'ISE', 'deviance'; default is "all"

### Value

<code>h</code>	cross-validation bandwidth for the chosen method; if no method was specified, then it is a list of three elements with entries corresponding to the estimated bandwidths on p-scale ( <code>h\$pscale</code> ), on eta-scale ( <code>h\$etascale</code> ) and for mean likelihood ( <code>h\$deviance</code> )
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bandwidth_plugin	<i>Plug in estimation of Bandwidth</i>
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---

**Description**

Calculates an estimate of the AMISE optimal bandwidth for the local polynomial estimator of a psychometric function.

**Usage**

```
bandwidth_plugin( r, m, x, link = c( "logit" ), guessing = 0, lapsing = 0, K = 2, p = 1, ker = c
```

**Arguments**

r	number of successes in points x
m	number of trials in points x
x	stimulus levels
link	link function; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2
p	order of the polynomial; default is 1
ker	kernel function for weights; default "dnorm"

**Value**

h	plug-in bandwidth (on eta-scale)
---	----------------------------------

**Examples**

```
data("01_Miranda")
h<-bandwidth_plugin( example01$r, example01$m, example01$x )
```

---

binomfit_lims	<i>Generalized linear model fit with guessing and lapsing rates</i>
---------------	---

---

**Description**

The function fits a binomial generalised liner model with fixed guessing and lapsing rates

**Usage**

```
binomfit_lims( r, m, x, p = 1, link = c( "logit" ), guessing = 0, lapsing = 0, K = 2 )
```

**Arguments**

r	number of successes in points x
m	number of trials in points x
x	stimulus levels
p	degree of the polynomial to be fitted on the linear scale; default is 1
link	link function; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2

**Value**

value	Object with 2 components: b: vector of estimated coefficients for the linear part fit: glm object to be used in evaluation of fitted values
-------	---

**Examples**

```
data( "01_Miranda" )
value <- binomfit_lims( example01$r, example01$m, example01$x )
```

binom\_g

*Psychometric function with guessing rate***Description**

THIS IS AN INTERNAL FUNCTION: USE BINOM\_LIMS FOR BEST RESULTS. Maximum likelihood estimates of the parameters of psychometric function with guessing rate (GLM). The estimated parameters for the linear part are in vector b and the estimated guessing rate is guess (GLM).

**Usage**

```
binom_g( r, m, x, link, p, K, initval )
```

**Arguments**

r	number of successes in points x
m	number of trials in points x
x	stimulus levels
link	link function
p	degree of the polynomial to be fitted on the linear scale
K	Power parameter in Weibull and reverse Weibull models
initval	initial value for guessing rate

**Value**

value	Object with 3 components: guessing: estimated guessing rate b: vector of estimated coefficients for the linear part fit: glm object to be used in evaluation of fitted values
-------	---

**Examples**

```
data( "01_Miranda" )
value <-binom_g( example01$r, example01$m, example01$x, "logit", 1, 2, 0.01 )
```

---

binom_gl	<i>Psychometric function with guessing and lapsing rates</i>
----------	--

---

**Description**

THIS IS AN INTERNAL FUNCTION: USE BINOM\_LIMS FOR BEST RESULTS. Maximum likelihood estimates of the parameters of psychometric function with guessing and lapsing rates (GLM) or only guessing rate. The estimated parameters for the linear part are in vector b and the estimated limits are in lims.

**Usage**

```
binom_gl( r, m, x, link, p, K, initval )
```

**Arguments**

r	number of successes in points x
m	number of trials in points x
x	stimulus levels
link	link function
p	degree of the polynomial to be fitted on the linear scale
K	Power parameter in Weibull and reverse Weibull models
initval	initial value for guessing and lapsing rates

**Value**

value	Object with 4 components: guessing: estimated guessing rate lapsing: estimated lapsing rate b: vector of estimated coefficients for the linear part fit: glm object to be used in evaluation of fitted values
-------	---

**Examples**

```
data( "01_Miranda" );
value <-binom_gl( example01$r, example01$m, example01$x, "logit", 1, 2, c( 0.01, 0.01 ) );
```

---

binom_l	<i>Psychometric function with lapsing rate</i>
---------	--

---

### Description

THIS IS AN INTERNAL FUNCTION: USE BINOM\_LIMS FOR BEST RESULTS. Maximum likelihood estimates of the parameters of psychometric function with lapsing rate (GLM). The estimated parameters for the linear part are in vector b and the estimated guessing rate is guess (GLM).

### Usage

```
binom_l( r, m, x, link, p, K, initval )
```

### Arguments

r	number of successes in points x
m	number of trials in points x
x	stimulus levels
link	link function
p	degree of the polynomial to be fitted on the linear scale
K	Power parameter in Weibull and reverse Weibull models
initval	initial value for lapsing rate

### Value

value	Object with 4 components: lapsing: estimated lapsing rate b: vector of estimated coefficients for the linear part fit: glm object to be used in evaluation of fitted values
-------	---

### Examples

```
data( "01_Miranda" )
value <-binom_l( example01$r, example01$m, example01$x, "logit", 1, 2, 0.01 )
```

---

binom_lims	<i>Psychometric function with guessing and lapsing rates</i>
------------	--

---

### Description

Maximum likelihood estimates of the parameters of psychometric function with guessing and lapsing rates (GLM) or only guessing rate. The estimated parameters for the linear part are in vector b and the estimated limits are in lims.

### Usage

```
binom_lims( r, m, x, gl = c( "both" ), link = c( "logit" ), p = 1, K = 2, initval = NULL )
```

**Arguments**

r	number of successes in points x
m	number of trials in points x
x	stimulus levels
gl	indicator, calculate only guessing if "guessing", only lapsing if "lapsing" and both guessing and lapsing if "both"; default is "both"
link	link function; default is "logit"
p	degree of the polynomial to be fitted on the linear scale; default is 1
K	Power parameter in Weibull and reverse Weibull models; default is 2
initval	initial value for guessing and lapsing; default is c( 0.01, 0.01) if guessing and lapsing rates are estimated, and 0.01 if only guessing or only lapsing rate is estimated

**Value**

value	The object returned by internal functions binom_g, binom_l, or binom_gl, depending on only guessing, only lapsing, or guessing and lapsing are estimated
-------	--

**Examples**

```
data( "01_Miranda" )
value <-binom_lims( example01$r, example01$m, example01$x )
```

---

binom_revweib	<i>Psychometric function fitting for reverse Weibull link function</i>
---------------	--

---

**Description**

Maximum likelihood estimates of the parameters of the reverser Weibull model(GLM). The estimated parameters for the linear part are in vector b and the estimated exponent is K.

**Usage**

```
binom_revweib( r, m, x, p = 1, initK = 2, guessing = 0, lapsing = 0 )
```

**Arguments**

r	number of successes in points x
m	number of trials in points x
x	stimulus levels
p	degree of the polynomial to be fitted on the linear scale; default is 1
initK	Power parameter in reverse Weibull model; default is 2
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0

**Value**

value                      Object with 3 components: b: vector of estimated coefficients for the linear part  
                              K: estimate of the power parameter in the reverse Weibull model fit: glm object  
                              to be used in evaluation of fitted values

**Examples**

```
data( "01_Miranda" )
value <- binom_revweib( example01$r, example01$m, example01$x )
```

binom\_weib

*Psychometric function fitting for Weibull link function***Description**

Maximum likelihood estimates of the parameters of the Weibull model (GLM). The estimated parameters for the linear part are in vector b and the estimated exponent is K.

**Usage**

```
binom_weib( r, m, x, p = 1, initK = 2, guessing = 0, lapsing = 0 )
```

**Arguments**

r                      number of successes in points x  
 m                      number of trials in points x  
 x                      stimulus levels  
 p                      degree of the polynomial to be fitted on the linear scale; default is 1  
 initK                  Power parameter in reverse Weibull model; default is 2  
 guessing              guessing rate; default is 0  
 lapsing                lapsing rate; default is 0

**Value**

value                      Object with 3 components: b: vector of estimated coefficients for the linear part  
                              K: estimate of the power parameter in the reverse Weibull model fit: glm object  
                              to be used in evaluation of fitted values

**Examples**

```
data( "01_Miranda" )
value <- binom_revweib( example01$r, example01$m, example01$x )
```

bootstrap\_ci\_sl

*Bootstrap estimate of confidence interval for slope estimation***Description**

Finds bootstrap estimate of a confidence interval at a significant level  $\alpha$  for the estimated slope for the local polynomial estimation of psychometric function (PF) with guessing and lapsing rates specified in `lims`. Confidence interval is based on bootstrap percentiles

**Usage**

```
bootstrap_ci_sl( TH, r, m, x, N, h0, alpha = 0.05, X = (max(x)-min(x))*(0:999)/999+min(x), link
```

**Arguments**

TH	required threshold level
r	number of successes in points x
m	number of trials in points x
x	stimulus levels
N	number of bootstrap replications
h0	pilot bandwidth; if not specified, then the scaled plug-in bandwidth is used
alpha	significance level of the confidence interval
X	set of value for which to calculate the estimates of PF for the threshold estimation; if not given 1000 equally spaced points from min to max of xdes are used
link	name of the link function to be used; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2
p	order of the polynomial; default is 1
ker	kernel function for weights; default "dnorm"
maxiter	maximum number of iterations in Fisher scoring; default is 50
tol	tolerance level at which to stop Fisher scoring; default is 1e-6

**Value**

value	Object with 2 components: <code>ci</code> : confidence interval based on bootstrap percentiles <code>sl0</code> : slope estimate
-------	--

**Examples**

```
data( "01_Miranda" )
bwd <- 0.2959
value <- bootstrap_ci_sl( 0.5, example01$r, example01$m, example01$x, 10, bwd )
```

bootstrap\_ci\_th

*Bootstrap estimate of confidence interval for threshold estimation***Description**

Finds bootstrap estimate of a confidence interval at a significant level  $\alpha$  for the estimated threshold for the local polynomial estimation of psychometric function (PF) with guessing and lapsing rates specified in `lims`. Confidence interval is based on bootstrap percentiles

**Usage**

```
bootstrap_ci_th( TH, r, m, x, N, h0, alpha = 0.05, X = (max(x)-min(x))*(0:999)/999+min(x), link
```

**Arguments**

TH	required threshold level
r	number of successes in points x
m	number of trials in points x
x	stimulus levels
N	number of bootstrap replications
h0	pilot bandwidth; if not specified, then the scaled plug-in bandwidth is used
alpha	significance level of the confidence interval
X	set of value for which to calculate the estimates of PF for the threshold estimation; if not given 1000 equally spaced points from min to max of xdes are used
link	name of the link function to be used; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2
p	order of the polynomial; default is 1
ker	kernel function for weights; default "dnorm"
maxiter	maximum number of iterations in Fisher scoring; default is 50
tol	tolerance level at which to stop Fisher scoring; default is 1e-6

**Value**

value	Object with 2 components: ci: confidence interval based on bootstrap percentiles th0: threshold estimate
-------	--

**Examples**

```
data( "01_Miranda" )
bwd <- 0.2959;
value <- bootstrap_ci_th( 0.5, example01$r, example01$m, example01$x, 10, bwd );
```



bootstrap\_sd\_sl

*Bootstrap estimate the standard deviation of slope estimation***Description**

Finds bootstrap estimate of the standard deviation of the estimated slope for the local polynomial estimation of psychometric function (PF) with guessing and lapsing rates as specified

**Usage**

```
bootstrap_sd_sl( TH, r, m, x, N, h0, X = (max(x)-min(x))*(0:999)/999+min(x), link = c( "logit" ) )
```

**Arguments**

TH	required threshold level
r	number of successes in points x
m	number of trials in points x
x	stimulus levels
N	number of bootstrap replications
h0	pilot bandwidth; if not specified, then the scaled plug-in bandwidth is used
X	set of value for which to calculate the estimates of PF for the threshold estimation; if not given 1000 equally spaced points from min to max of xdes are used
link	name of the link function to be used; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2
p	order of the polynomial; default is 1
ker	kernel function for weights; default "dnorm"
maxiter	maximum number of iterations in Fisher scoring; default is 50
tol	tolerance level at which to stop Fisher scoring; default is 1e-6

**Value**

value	Object with 2 components: sd: bootstrap estimate of the standard deviation of the slope estimate sl0: slope estimate
-------	--

**Examples**

```
data( "01_Miranda" )
bwd <- 0.2959
value <- bootstrap_sd_sl( 0.5, example01$r, example01$m, example01$x, 10, bwd )
```

bootstrap\_sd\_th

*Bootstrap estimate the standard deviation of threshold estimation***Description**

Finds bootstrap estimate of the standard deviation of the estimated threshold for the local polynomial estimation of psychometric function (PF) with guessing and lapsing rates as specified

**Usage**

```
bootstrap_sd_th( TH, r, m, x, N, h0, X = (max(x)-min(x))*(0:999)/999+min(x), link = c( "logit" ) )
```

**Arguments**

TH	required threshold level
r	number of successes in points x
m	number of trials in points x
x	stimulus levels
N	number of bootstrap replications
h0	pilot bandwidth; if not specified, then the scaled plug-in bandwidth is used
X	set of value for which to calculate the estimates of PF for the threshold estimation; if not given 1000 equally spaced points from min to max of xdes are used
link	name of the link function to be used; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2
p	order of the polynomial; default is 1
ker	kernel function for weights; default "dnorm"
maxiter	maximum number of iterations in Fisher scoring; default is 50
tol	tolerance level at which to stop Fisher scoring; default is 1e-6

**Value**

value      Object with 2 components: sd: bootstrap estimate of the standard deviation of the threshold estimate th0: threshold estimate

**Examples**

```
data( "01_Miranda" )
bwd <- 0.2959
value <- bootstrap_sd_th( 0.5, example01$r, example01$m, example01$x, 10, bwd )
```

---

checkinput	<i>Check input parameters of modelfree functions</i>
------------	--

---

**Description**

THIS IS AN INTERNAL FUNCTION ONLY. Pool of routines that check robustness of input of parameters passed to the other functions of modelfree package.

**Usage**

```
checkinput(type,x)
```

**Arguments**

type	Type of checking
x	Input data to be checked

**Value**

None. Only stops execution if data is not consistent.

---

comploglog_link	<i>Complementary log-log link function with guessing and lapsing rates</i>
-----------------	--

---

**Description**

Complementary loglog link for use with GLM functions. The guessing and lapsing rate are fixed to values given in lms, hence link is a function of only one variable

**Usage**

```
comploglog_link( guessing = 0, lapsing = 0 )
```

**Arguments**

guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0

**Value**

link	Complementary log-log link for use in all GLM functions
------	---

**Examples**

```

data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m ,m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-comploglog_link( 0.1, 0.1 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )

```

---

comploglog\_link\_private

*Complementary log-log link function with guessing and lapsing rates*

---

**Description**

THIS IS AN INTERNAL FUNCTION: USE COMPLOGLOG\_LINK FOR BEST RESULTS. Complementary loglog link for use with GLM functions. The guessing and lapsing rate are fixed to values given in lims, hence link is a function of only one variable

**Usage**

```
comploglog_link_private( guessing, lapsing )
```

**Arguments**

guessing	guessing rate
lapsing	lapsing rate

**Value**

link	Complementary log-log link for use in all GLM functions
------	---

**Examples**

```

data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m ,m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-comploglog_link_private( 0.1, 0.1 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )

```

---

deviance2	<i>Deviance between data and fitted function</i>
-----------	--

---

**Description**

Calculates deviance for data Y and fitted values of the psychometric function f

**Usage**

```
deviance2(r, m, pfit)
```

**Arguments**

r	number of successes in points x
m	number of trials in points x
pfit	fitted values

**Value**

D	Deviance
---	----------

**Examples**

```
data( "01_Miranda" )
h = 0.2959
fit <- locglmfit( example01$x, example01$r, example01$m, example01$x, h )
Dev <- deviance2( example01$r, example01$m, fit$fitval )
```

---

locglmfit	<i>Local generalized linear fitting</i>
-----------	---

---

**Description**

Local polynomial estimator for the psychometric function (PF) and eta function (PF transformed by link) for binomial data; also returns the Hat matrix. Actual calculations are done in LOCGLMFIT\_PRIVATE or LOCGLMFIT\_SPARSE\_PRIVATE depending on the size of the data set. Here the data are split into several parts to speed up the calculations.

**Usage**

```
locglmfit( xfit, r, m, x, h, returnH = FALSE, link = c( "logit" ), guessing = 0, lapsing = 0, K
```

**Arguments**

xfit	points in which to calculate the estimate
r	number of successes in points x
m	number of trials in points x
x	stimulus values
h	bandwidths
returnH	Boolean; Return or not the hat matrix H? default is TRUE
link	name of the link function to be used; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2
p	degree of the polynomial; default p = 1
ker	kernel function for weights; default "dnorm"
maxiter	maximum number of iterations in Fisher scoring; default is 50
tol	tolerance level at which to stop Fisher scoring; default is 1e-6

**Value**

value	Object with 2 or 3 components: pfit: value of the local polynomial estimate at points xfit etafit: estimate of eta (link of pfit) H: hat matrix (OPTIONAL)
-------	--

**Examples**

```
data( "01_Miranda" )
xnew = 1.2 * (0:99)/99+0.1
h <- 0.2959
fit <- locglmfit( xnew, example01$r, example01$m, example01$x, h )
```

---

locglmfit_private	<i>Local generalized linear fitting with usual (non-sparse) matrices</i>
-------------------	--

---

**Description**

THIS IS AN INTERNAL FUNCTION: USE LOCGLMFIT FOR BEST RESULTS. Fisher scoring method for local polynomial estimator of a psychometric function (PF).

**Usage**

```
locglmfit_private( xfit, r, m, x, h, returnH, link, guessing, lapsing, K, p, ker, maxiter, tol )
```

**Arguments**

xfit	points in which to calculate the estimate
r	number of successes in points x
m	number of trials in points x
x	stimulus values
h	bandwidths
returnH	Boolean; Return or not the hat matrix H? default is TRUE
link	name of the link function to be used; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2
p	degree of the polynomial; default p = 1
ker	kernel function for weights; default "dnorm"
maxiter	maximum number of iterations in Fisher scoring; default is 50
tol	tolerance level at which to stop Fisher scoring; default is 1e-6

**Value**

value	Object with 2 or 3 components: pfit: value of the local polynomial estimate at points xfit etafit: estimate of eta (link of pfit) H: hat matrix (OPTIONAL)
-------	--

**Examples**

```
data( "01_Miranda" )
xnew = 1.2 * (0:99)/99+0.1
h <- 0.2959
fit <- locglmfit_private( xnew, example01$r, example01$m, example01$x, h, FALSE, "logit_link", 0, 0, 2, 1
```

---

locglmfit\_sparse\_private

*Local generalized linear fitting with sparse matrices*


---

**Description**

THIS IS AN INTERNAL FUNCTION: USE LOCGLMFIT FOR BEST RESULTS. Fisher scoring method for local polynomial estimator of a psychometric function (PF).

**Usage**

```
locglmfit_sparse_private( xfit, r, m, x, h, returnH, link, guessing, lapsing, K, p, ker, maxiter
```

**Arguments**

xfit	points in which to calculate the estimate
r	number of successes in points x
m	number of trials in points x
x	stimulus values
h	bandwidths
returnH	Boolean; Return or not the hat matrix H? default is TRUE
link	name of the link function to be used; default is "logit"
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0
K	power parameter for Weibull and reverse Weibull link; default is 2
p	degree of the polynomial; default p = 1
ker	kernel function for weights; default "dnorm"
maxiter	maximum number of iterations in Fisher scoring; default is 50
tol	tolerance level at which to stop Fisher scoring; default is 1e-6

**Value**

value	Object with 2 or 3 components: pfit: value of the local polynomial estimate at points xfit etafit: estimate of eta (link of pfit) H: hat matrix (OPTIONAL)
-------	--

**Examples**

```
data( "01_Miranda" )
xnew = 1.2 * (0:99)/99+0.1
h <- 0.2959
fit <- locglmfit_sparse_private( xnew, example01$r, example01$m, example01$x, h, FALSE, "logit_link", 0,
```

logit\_link

*Logit link function with guessing and lapsing rates***Description**

Logit link for use with GLM functions. The guessing and lapsing rate are fixed to values given in `lims`, hence `link` is a function of only one variable

**Usage**

```
logit_link( guessing = 0, lapsing = 0 )
```

**Arguments**

guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0

**Value**

link	Logit link for use in all GLM functions
------	---



**Examples**

```

data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m , m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-logit_link( 0.1, 0.1 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )

```

---

logit_link_private	<i>Logit link function with guessing and lapsing rates</i>
--------------------	--

---

**Description**

THIS IS AN INTERNAL FUNCTION: USE LOGIT\_LINK FOR BEST RESULTS. Logit link for use with GLM functions. The guessing and lapsing rate are fixed to values given in lims, hence link is a function of only one variable

**Usage**

```
logit_link_private( guessing, lapsing )
```

**Arguments**

guessing	guessing rate
lapsing	lapsing rate

**Value**

link	Logit link for use in all GLM functions
------	---

**Examples**

```

data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m , m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-logit_link_private( 0.1, 0.1 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )

```

---

loglog_link	<i>Log-log link function with guessing and lapsing rates</i>
-------------	--

---

**Description**

Log-log link for use with GLM functions. The guessing and lapsing rate are fixed to values given in `lims`, hence `link` is a function of only one variable

**Usage**

```
loglog_link( guessing = 0, lapsing = 0 )
```

**Arguments**

<code>guessing</code>	guessing rate; default is 0
<code>lapsing</code>	lapsing rate; default is 0

**Value**

<code>link</code>	Log-log link for use in all GLM functions
-------------------	---

**Examples**

```
data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m , m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-loglog_link( 0.1, 0.1 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )
```

---

loglog_link_private	<i>Log-log link function with guessing and lapsing rates</i>
---------------------	--

---

**Description**

THIS IS AN INTERNAL FUNCTION: USE `CLOGLOG_LINK` FOR BEST RESULTS. Log-log link for use with GLM functions. The guessing and lapsing rate are fixed to values given in `lims`, hence `link` is a function of only one variable

**Usage**

```
loglog_link_private( guessing, lapsing )
```

**Arguments**

<code>guessing</code>	guessing rate
<code>lapsing</code>	lapsing rate

**Value**

link                      Log-log link for use in all GLM functions

**Examples**

```
data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m , m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-loglog_link_private( 0.1, 0.1 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )
```

---

probit_link	<i>Probit link function with guessing and lapsing rates</i>
-------------	---

---

**Description**

Probit link for use with GLM functions. The guessing and lapsing rate are fixed to values given in `lims`, hence link is a function of only one variable

**Usage**

```
probit_link( guessing = 0, lapsing = 0 )
```

**Arguments**

guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0

**Value**

link                      Probit link for use in all GLM functions

**Examples**

```
data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m , m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-probit_link( 0.1, 0.1 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )
```

---

probit_link_private	<i>Probit link function with guessing and lapsing rates</i>
---------------------	---

---

### Description

THIS IS AN INTERNAL FUNCTION: USE PROBIT\_LINK FOR BEST RESULTS. Probit link for use with GLM functions. The guessing and lapsing rate are fixed to values given in lms, hence link is a function of only one variable

### Usage

```
probit_link_private( guessing, lapsing )
```

### Arguments

guessing	guessing rate
lapsing	lapsing rate

### Value

link	Probit link for use in all GLM functions
------	--

### Examples

```
data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m , m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-probit_link_private( 0.1, 0.1 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )
```

---

revweibull_link	<i>Reverse Weibull link function with guessing and lapsing rates</i>
-----------------	--

---

### Description

Reverse Weibull link for use with GLM functions. The guessing rate and lapsing rate are fixed, and power parameter is set to be equal K, hence link is a function of only one variable

### Usage

```
revweibull_link( K, guessing = 0, lapsing = 0 )
```

### Arguments

K	power parameter for reverse Weibull link function
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0

**Value**

link	Reverse Weibull link for use in all GLM functions
------	---

**Examples**

```
data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m ,m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-revweibull_link( 20 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )
```

---

revweibull\_link\_private

*Reverse Weibull link function with guessing and lapsing rates*


---

**Description**

THIS IS AN INTERNAL FUNCTION: USE REVWEIBULL\_LINK FOR BEST RESULTS. Reverse Weibull link for use with GLM functions. The guessing rate and lapsing rate are fixed, and power parameter is set to be equal K, hence link is a function of only one variable

**Usage**

```
revweibull_link_private( K, guessing, lapsing )
```

**Arguments**

K	power parameter for reverse Weibull link function
guessing	guessing rate
lapsing	lapsing rate

**Value**

link	Reverse Weibull link for use in all GLM functions
------	---

**Examples**

```
data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m ,m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-revweibull_link_private( 20, 0, 0 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )
```

---

threshold_slope	<i>Threshold and slope of estimated psychometric function</i>
-----------------	---

---

### Description

Finds the approximate value of  $x$  ( $=x_{th}$ ) for which the value of the psychometric function is equal thresh and the approximate value of slope in  $x_{th}$

### Usage

```
threshold_slope( pfit, xfit, thresh = 0.5 )
```

### Arguments

pfit	estimated values of the psychometric function
xfit	stimulus levels in which the function was estimated
thresh	value for which to estimate threshold; default is 0.5

### Value

value	Object with 2 elements: $x_{th}$ : estimated threshold slope: estimated value of slope, i.e. derivative of pfit at $x_{th}$
-------	---

### Examples

```
data( "01_Miranda" )
xnew = 1.2 * (0:999)/999+0.1
h = 0.2959
fit <- locglmfit( xnew, example01$r, example01$m, example01$x, h )
value <- threshold_slope( fit$pfit , xnew )
```

---

weibull_link	<i>Weibull link function with guessing and lapsing rates</i>
--------------	--

---

### Description

Weibull link for use with GLM functions. The guessing rate and lapsing rate are fixed, and power parameter is set to be equal K, hence link is a function of only one variable

### Usage

```
weibull_link( K, guessing = 0, lapsing = 0 )
```

### Arguments

K	power parameter for reverse Weibull link function
guessing	guessing rate; default is 0
lapsing	lapsing rate; default is 0

**Value**

link	Weibull link for use in all GLM functions
------	---

**Examples**

```
data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m ,m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-weibull_link( 20 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )
```

---

weibull_link_private	<i>Weibull link function with guessing and lapsing rates</i>
----------------------	--

---

**Description**

THIS IS AN INTERNAL FUNCTION: USE WEIBULL\_LINK FOR BEST RESULTS. Weibull link for use with GLM functions. The guessing rate and lapsing rate are fixed, and power parameter is set to be equal K, hence link is a function of only one variable

**Usage**

```
weibull_link_private( K, guessing, lapsing )
```

**Arguments**

K	power parameter for reverse Weibull link function
guessing	guessing rate
lapsing	lapsing rate

**Value**

link	Weibull link for use in all GLM functions
------	---

**Examples**

```
data( "01_Miranda" )
x <- example01$x
r <- example01$r
m <- example01$m
glmdata <- data.frame( cbind( r/m ,m , x ) )
names( glmdata ) <- c( "resp", "m", "x" )
glmformula <- c( "resp ~ x" )
userlink<-weibull_link_private( 20, 0, 0 )
fit <- glm( glmformula, data = glmdata, weights = m, family = binomial( userlink ) )
```

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